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SEQUENCE LISTING

<110> Moon, Chulso
Mao, Li

<120> DAP-Kinase and HOXA9, Two Human Genes Associated with Genesis,
Progression, and Aggressiveness of Non-Small Cell Lung Cancer

<130> 10620-1U1

<140> 10/045,400

<141> 2001-11-29

<150> US 60/250,083

<151> 2000-11-29

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Glu Cys Leu Ala Glu His Gly Ala Asp Leu Asn Ala Cys Asp Lys Asp
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Gly His Ile Ala Leu His Leu Ala Val Arg Arg Cys Gln Met Glu Val
 545 550 555 560

Ile Lys Thr Leu Leu Ser Gln Gly Cys Phe Val Asp Tyr Gln Asp Arg
 565 570 575

His Gly Asn Thr Pro Leu His Val Ala Cys Lys Asp Gly Asn Met Pro
 580 585 590

Ile Val Val Ala Leu Cys Glu Ala Asn Cys Asn Leu Asp Ile Ser Asn
 595 600 605

Lys Tyr Gly Arg Thr Pro Leu His Leu Ala Ala Asn Asn Gly Ile Leu
 610 615 620

Asp Val Val Arg Tyr Leu Cys Leu Met Gly Ala Ser Val Glu Ala Leu
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Thr Thr Asp Gly Lys Thr Ala Glu Asp Leu Ala Arg Ser Glu Gln His
 645 650 655

Glu His Val Ala Gly Leu Leu Ala Arg Leu Arg Lys Asp Thr His Arg
 660 665 670

Gly Leu Phe Ile Gln Gln Leu Arg Pro Thr Gln Asn Leu Gln Pro Arg
 675 680 685

Ile Lys Leu Lys Leu Phe Gly His Ser Gly Ser Gly Lys Thr Thr Leu
 690 695 700

Val Glu Ser Leu Lys Cys Gly Leu Leu Arg Ser Phe Phe Arg Arg Arg
 705 710 715 720

Arg Pro Arg Leu Ser Ser Thr Asn Ser Ser Arg Phe Pro Pro Ser Pro
 725 730 735

Leu Ala Ser Lys Pro Thr Val Ser Val Ser Ile Asn Asn Leu Tyr Pro
 740 745 750

Gly Cys Glu Asn Val Ser Val Arg Ser Arg Ser Met Met Phe Glu Pro
 755 760 765

Gly Leu Thr Lys Gly Met Leu Glu Val Phe Val Ala Pro Thr His His
 770 775 780

Pro His Cys Ser Ala Asp Asp Gln Ser Thr Lys Ala Ile Asp Ile Gln
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Asn Ala Tyr Leu Asn Gly Val Gly Asp Phe Ser Val Trp Glu Phe Ser
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Gly Asn Pro Val Tyr Phe Cys Cys Tyr Asp Tyr Phe Ala Ala Asn Asp

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Pro Thr Ser Ile His Val Val Val Phe Ser Leu Glu Glu Pro Tyr Glu		
835	840	845
Ile Gln Leu Asn Pro Val Ile Phe Trp Leu Ser Phe Leu Lys Ser Leu		
850	855	860
Val Pro Val Glu Glu Pro Ile Ala Phe Gly Gly Lys Leu Lys Asn Pro		
865	870	875 880
Leu Gln Val Val Leu Val Ala Thr His Ala Asp Ile Met Asn Val Pro		
	885	890 895
Arg Pro Ala Gly Gly Glu Phe Gly Tyr Asp Lys Asp Thr Ser Leu Leu		
	900	905 910
Lys Glu Ile Arg Asn Arg Phe Gly Asn Asp Leu His Ile Ser Asn Lys		
	915	920 925
Leu Phe Val Leu Asp Ala Gly Ala Ser Gly Ser Lys Asp Met Lys Val		
	930	935 940
Leu Arg Asn His Leu Gln Glu Ile Arg Ser Gln Ile Val Ser Val Cys		
945	950	955 960
Pro Pro Met Thr His Leu Cys Glu Lys Ile Ile Ser Thr Leu Pro Ser		
	965	970 975
Trp Arg Lys Leu Asn Gly Pro Asn Gln Leu Met Ser Leu Gln Gln Phe		
	980	985 990
Val Tyr Asp Val Gln Asp Gln Leu Asn Pro Leu Ala Ser Glu Glu Asp		
	995	1000 1005
Leu Arg Arg Ile Ala Gln Gln Leu His Ser Thr Gly Glu Ile Asn		
1010	1015	1020
Ile Met Gln Ser Glu Thr Val Gln Asp Val Leu Leu Leu Asp Pro		
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Arg Trp Leu Cys Thr Asn Val Leu Gly Lys Leu Leu Ser Val Glu		
1040	1045	1050

Thr	Pro	Arg	Ala	Leu	His	His	Tyr	Arg	Gly	Arg	Tyr	Thr	Val	Glu
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	1070					1075					1080			
Gln	Ile	Leu	Asp	Ala	Met	Asp	Ile	Cys	Ala	Arg	Asp	Leu	Ser	Ser
	1085					1090					1095			
Gly	Thr	Met	Val	Asp	Val	Pro	Ala	Leu	Ile	Lys	Thr	Asp	Asn	Leu
	1100					1105					1110			
His	Arg	Ser	Trp	Ala	Asp	Glu	Glu	Asp	Glu	Val	Met	Val	Tyr	Gly
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	1130					1135					1140			
Gly	Ile	Phe	His	Lys	Val	Gln	Val	Asn	Leu	Cys	Arg	Trp	Ile	His
	1145					1150					1155			
Gln	Gln	Ser	Thr	Glu	Gly	Asp	Ala	Asp	Ile	Arg	Leu	Trp	Val	Asn
	1160					1165					1170			
Gly	Cys	Lys	Leu	Ala	Asn	Arg	Gly	Ala	Glu	Leu	Leu	Val	Leu	Leu
	1175					1180					1185			
Val	Asn	His	Gly	Gln	Gly	Ile	Glu	Val	Gln	Val	Arg	Gly	Leu	Glu
	1190					1195					1200			
Thr	Glu	Lys	Ile	Lys	Cys	Cys	Leu	Leu	Leu	Asp	Ser	Val	Cys	Ser
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Thr	Ile	Glu	Asn	Val	Met	Ala	Thr	Thr	Leu	Pro	Gly	Leu	Leu	Thr
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Ser Phe Ser Ser Ile Met Cys Phe Gly Cys His Asp Val Tyr Ser
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Gln Ala Ser Leu Gly Met Asp Ile His Ala Ser Asp Leu Asn Leu
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Leu Thr Arg Arg Lys Leu Ser Arg Leu Leu Asp Pro Pro Asp Pro
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Leu Gly Lys Asp Trp Cys Leu Leu Ala Met Asn Leu Gly Leu Pro
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Asp Leu Val Ala Lys Tyr Asn Thr Asn Asn Gly Ala Pro Lys Asp
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Phe Leu Pro Ser Pro Leu His Ala Leu Leu Arg Glu Trp Thr Thr
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Tyr Pro Glu Ser Thr Val Gly Thr Leu Met Ser Lys Leu Arg Glu
 1370 1375 1380

Leu Gly Arg Arg Asp Ala Ala Asp Leu Leu Leu Lys Ala Ser Ser
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Val Phe Lys Ile Asn Leu Asp Gly Asn Gly Gln Glu Ala Tyr Ala
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Asp Leu Gly Ser Phe Leu Leu Leu Gly Ser Thr Phe Leu Ser Thr Gly	
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Thr Thr Leu Pro Phe Ile Thr Ser Val Glu Ile Val Ser Arg Tyr Leu	
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Cys Ala Arg Gly Ser Gly Arg Ala Gly His His Gly Pro Gly Arg Ala	
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agg cct gcg gtg gca acc tcg gct ttc cct gct cag gag cct cgt gtc	288
Arg Pro Ala Val Ala Thr Ser Ala Phe Pro Ala Gln Glu Pro Arg Val	
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Phe Leu Arg Ser Ala Leu Pro Ala Gly Arg Leu Ser Pro Ser Thr Thr	
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His Leu His Leu Val Thr Ala Asp Asn Pro Ala Ala Asn Trp Leu His	
115 120 125	
gcg cgc tcc act cgg aaa aag cgg tgc ccc tat aca aaa cac cag acc	432
Ala Arg Ser Thr Arg Lys Lys Arg Cys Pro Tyr Thr Lys His Gln Thr	
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Leu Glu Leu Glu Lys Glu Phe Leu Phe Asn Met Tyr Leu Thr Arg Asp	
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35 40 45

Thr Thr Leu Pro Phe Ile Thr Ser Val Glu Ile Val Ser Arg Tyr Leu
50 55 60

Cys Ala Arg Gly Ser Gly Arg Ala Gly His His Gly Pro Gly Arg Ala
65 70 75 80

Arg Pro Ala Val Ala Thr Ser Ala Phe Pro Ala Gln Glu Pro Arg Val
85 90 95

Phe Leu Arg Ser Ala Leu Pro Ala Gly Arg Leu Ser Pro Ser Thr Thr
100 105 110

His Leu His Leu Val Thr Ala Asp Asn Pro Ala Ala Asn Trp Leu His
115 120 125

Ala Arg Ser Thr Arg Lys Lys Arg Cys Pro Tyr Thr Lys His Gln Thr
130 135 140

Leu Glu Leu Glu Lys Glu Phe Leu Phe Asn Met Tyr Leu Thr Arg Asp
145 150 155 160

Arg Arg Tyr Glu Val Ala Arg Leu Leu Asn Leu Thr Glu Arg Gln Val
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Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Met Lys Lys Ile Asn Lys
180 185 190

Asp Arg Ala Lys Asp Glu
195